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Amendments to the Claims:

This listing of claims replaces all prior versions and listings of claims in the application:

Listing of Claims:

1. (Original) A polypeptide having the ability to bind CEA comprising the amino acid sequence:

wherein:

X₄ is Asn, Glu, Asp, or Met;

X₅ is Leu, Phe, Tyr, Trp, Val, Met, Ile, or Asn;

X₆ is Phe, Leu, Asp, Glu, Ala, Ile, Lys, Asn, Ser, Val, Trp, Tyr, Gly, or Thr;

X₇ is Lys, Phe, Asp, Gly, Leu, Asn, Trp, Ala, Gln, or Thr;

X₈ is Asn, Pro, Phe, Gly, Asp, Ala, Ser, Glu, Gln, Trp, His, Arg, Met, Val, or Leu;

X₉ is Gln, Lys, Leu, or Gly;

 X_{10} is Trp, Ala, or Tyr; and

X₁₁ is Phe, Thr, Met, Ser, Ala, Asn, Val, His, Ile, Pro, Trp, Tyr, Gly, Leu, or Glu.

2. (Original) A polypeptide having the ability to bind CEA comprising the amino acid sequence:

$$X_1$$
- X_2 - X_3 -Cys- X_4 - X_5 - X_6 - X_7 - X_8 - X_9 - X_{10} - X_{11} -Cys- X_{12} - X_{13} - X_{14} , (SEQ ID NO:111), wherein:

 X_1 is Asp, Asn, Ala, or Ile;

 X_2 is Trp;

X₃ is Val, Ile, Met, Tyr, Phe, Pro, or Asp;

X₄ is Asn, Glu, Asp, or Met;

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X₅ is Leu, Phe, Tyr, Trp, Val, Met, Ile, or Asn;

X₆ is Phe, Leu, Asp, Glu, Ala, Ile, Lys, Asn, Ser, Val, Trp, Tyr, Gly, or Thr;

X₇ is Lys, Phe, Asp, Gly, Leu, Asn, Trp, Ala, Gln, or Thr;

X₈ is Asn, Pro, Phe, Gly, Asp, Ala, Ser, Glu, Gin, Trp, His, Arg, Met, Val, or Leu;

X₉ is Gln, Lys, Leu, or Gly;

 X_{10} is Trp, Ala, or Tyr; and

X₁₁ is Phe, Thr, Met, Ser, Ala, Asn, Val, His, Ile, Pro, Trp, Tyr, Gly, Leu, or Glu.

X₁₂ is Asn, Asp, Glu, Pro, Gln, Ser, Phe, or Val;

X₁₃ is Val, Leu, Ile, Pro, Ala, Gln, Ser, Met, Glu, Thr, Lys, Trp, or Arg; and

X₁₄ is Leu, Met, Val, Tyr, Ala, Ile, Trp, His, Pro, Gln, Glu, Phe, Lys, Arg, or Ser.

3. (Previously presented) A polypeptide having the ability to bind CEA comprising the amino acid sequence:

Cys-
$$X_4$$
- X_5 - X_6 - X_7 - X_8 - X_9 - X_{10} - X_{11} -Cys, (SEQ ID NO:3)

wherein:

X₄ is Asn, Glu, or Met;

X₅ is Asn, Leu, Met or Phe;

 X_6 is Asp, Gly, Ile, Lys Phe or Thr;

 X_7 is Ala, Gln, Gly, Lys or Thr;

X₈ is Arg, Asn, Asp, Glu or Gly;

X₉ is Gln, Gly or Leu;

 X_{10} is Ala, Trp or Tyr;

 X_{11} is Ala, Gly, His, Phe, Thr or Val.

4. (Original) The polypeptide according to claim 3, wherein:

X₄ is Glu;

X₅ is Asn, Leu, Met or Phe;

X₆ is Asp, Gly, Ile, Lys Phe or Thr;

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X₇ is Lys;

X₈ is Arg, Asn, Asp, Glu or Gly;

X₉ is Gln;

 X_{10} is Trp;

 X_{11} is Ala, Gly, His, Phe, Thr or Val.

5. (Original) The polypeptide according to claim 3, comprising the amino acid sequence:

$$X_1-X_2-X_3-Cys-X_4-X_5-X_6-X_7-X_8-X_9-X_{10}-X_{11}-Cys-X_{12}-X_{13}-X_{14}$$
, (SEQ ID NO:1),

wherein:

 X_1 is Asn or Asp;

 X_2 is Trp;

X₃ is Asp, Phe or Val;

X₄ is Asn, Glu or Met;

X₅ is Asn, Leu, Met or Phe;

X₆ is Asp, Gly, Ile, Lys, Phe or Thr;

 X_7 is Ala, Gln, Gly, Lys or Thr;

X₈ is Arg, Asn, Asp, Glu or Gly;

X₉ is Gln, Gly or Leu;

 X_{10} is Ala, Trp or Tyr;

X₁₁ is Ala, Gly, His, Phe, Thr or Val;

 X_{12} is Asn, Gln, Phe, Ser or Val;

 X_{13} is Arg, Leu, Pro or Ser; and

 X_{14} is Leu, Ser, Trp or Tyr.

6. (Previously presented) The polypeptide according to claim 5, having the amino acid sequence:

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 X_1 -Trp-Val-Cys-Glu- X_5 - X_6 -Lys- X_8 -Gln-Trp- X_{11} -Cys-Asn- X_{13} - X_{14} (SEQ ID NO:2), wherein:

 X_1 is Asn or Asp;

X₅ is Asn, Leu, Met or Phe;

X₆ is Asp, Gly, Ile, Lys, Phe or Thr;

X₈ is Arg, Asn, Asp, Glu or Gly;

X₁₁ is Ala, Gly, His, Phe, Thr or Val;

 X_{13} is Arg, Leu, Pro or Ser; and

 X_{14} is Leu or Tyr.

7. (Original) The polypeptide according to claim 5, comprising an amino acid sequence selected from the group consisting of:

Asn-Trp-Val-Cys-Asn-Leu-Phe-Lys-Asn-Gln-Trp-Phe-Cys-Asn-Ser-Tyr; (SEQ ID NO:4);

Asp-Trp-Val-Cys-Glu-Asn-Lys-Lys-Asp-Gln-Trp-Thr-Cys-Asn-Leu-Leu; (SEQ ID NO:5);

Asn-Trp-Asp-Cys-Met-Phe-Gly-Ala-Glu-Gly-Trp-Ala-Cys-Ser-Pro-Trp; (SEQ ID NO:6);

Asp-Trp-Val-Cys-Glu-Lys-Thr-Thr-Gly-Gly-Tyr-Val-Cys-Gln-Pro-Leu; (SEQ ID NO:7);

Asn-Trp-Phe-Cys-Glu-Met-Ile-Gly-Arg-Gln-Trp-Gly-Cys-Val-Pro-Ser; (SEQ ID NO:8); and

Asp-Trp-Val-Cys-Asn-Phe-Asp-Gln-Gly-Leu-Ala-His-Cys-Phe-Pro-Ser. (SEQ ID NO:9).

8. (Original) A polypeptide having the ability to bind CEA comprising the amino acid sequence:

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 $X_1-X_2-X_3-Cys-X_4-X_5-X_6-X_7-X_8-X_9-X_{10}-X_{11}-Cys-X_{12}-X_{13}-X_{14}$, (SEQ ID NO:1), wherein:

 X_1 is Asp, Asn, Ala, or Ile;

 X_2 is Trp;

X₃ is Val, Ile, Met, Tyr, Phe, Pro, or Asp;

X₄ is Asn, Glu, or Asp;

X₅ is Leu, Phe, Tyr, Trp, Val, Met, Ile, or Asn;

X₆ is Phe, Leu, Asp, Glu, Ala, Ile, Lys, Asn, Ser, Val, Trp, or Tyr;

X₇ is Lys, Phe, Asp, Gly, Leu, Asn, or Trp;

X₈ is Asn, Pro, Phe, Gly, Asp, Ala, Ser, Glu, Gln, or Trp;

X₉ is Gln, or Lys;

 X_{10} is Trp;

X₁₁ is Phe, Thr, Met, Ser, Ala, Asn, Val, His, Ile, Pro, Trp, or Tyr;

X₁₂ is Asn, Asp, Glu, Pro, Gln, or Ser;

X₁₃ is Val, Leu, Ile, Pro, Ala, Gln, Ser, Met, Glu, Thr, Lys, or Trp; and

X₁₄ is Leu, Met, Val, Tyr, Ala, Ile, Trp, His, Pro, Gln, Glu, Phe, Lys, or Arg.

9. (Previously presented) The polypeptide of Claim 1, wherein:

X₄ is Asn, or Glu;

X₅ is Leu, Phe, Tyr, Trp, or Ile;

X₆ is Phe, Leu, Asp, Glu, Ile, Ser, Val, or Gly;

 X_7 is Lys;

X₈ is Asn, Pro, Gly, Asp, Ala, Ser, His, Met, Val, or Leu;

X₉ is Gln;

 X_{10} is Trp;

X₁₁ is Phe, Thr, Ser, Ala, Asn, Val, His, Ile, Trp, Tyr, Leu, or Glu.

10. (Previously presented) The polypeptide of Claim 2, wherein:

 X_1 is Asp, or Asn;

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 X_2 is Trp;

X₃ is Val, Ile, or Met;

X₄ is Asn, or Glu;

X₅ is Leu, Phe, Tyr, Trp, or Ile;

X₆ is Phe, Leu, Asp, Glu, Ile, Ser, Val, or Gly;

X₇ is Lys;

X₈ is Asn, Pro, Gly, Asp, Ala, Ser, His, Met, Val, or Leu;

X₉ is Gln;

 X_{10} is Trp;

X₁₁ is Phe, Thr, Ser, Ala, Asn, Val, His, Ile, Trp, Tyr, Leu, or Glu;

 X_{12} is Asn, or Asp;

X₁₃ is Val, Leu, Ile, Pro, Ala, Gln, Ser, or Met; and

X₁₄ is Leu, Met, Val, Tyr, Trp, His, Gln, Arg, or Ser.

- 11. (Previously presented) The polypeptide according to Claim 2, comprising an amino acid sequence selected from the group consisting of SEQ ID NOs: 37-109 and 113-151.
- 12. (Original) The polypeptide according to Claim 1, 2, 3, 5, 8, 9, or 10, wherein said polypeptide binds to CEA but does not bind to NCA.
- 13. (Previously presented) The polypeptide according to claim 1, 2, 3, 5, 8, 9, or 10, wherein said polypeptide has a K_d for CEA which is less than 7 μ M.

14. -30. (Cancel)